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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2007; month=11; day=25; hr=15; min=27; sec=47; ms=845;
]

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Reviewer Comments:

<210> 2
<211> 372
<212> PRT
<213> Chemically synthesized

The above <213> response is invalid: per Sequence Rules, the only valid responses are: the Genus species of the organism, "Artificial Sequence," or "Unknown." "Artificial Sequence" and "Unknown" require explanation in the <220>-<223> section.

Application No: 10565751 Version No: 1.0

Input Set:

Output Set:

Started: 2007-11-02 19:28:57.317
Finished: 2007-11-02 19:28:57.902
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 585 ms
Total Warnings: 6
Total Errors: 0
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
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SEQUENCE LISTING

<110> INSTITUTE OF MEDICINAL BIOTECHNOLOGY, CHINESE ACADEMY OF MEDICAL SCIENCES

<120> INTENSIFIED FUSION PROTEIN FV-LDP-AW HAVING ANGIOGENESIS INHIBITING AND ANTITUMOR ACTIVITY AND THE USE THEREOF

<130> 57000/C306

<140> 10565751

<141> 2007-11-02

<160> 6

<170> PatentIn version 3.4

<210> 1

<211> 1119

<212> DNA

<213> Artificial sequence

<220>

<223> Chemically synthesized

<400> 1

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acgcctgaac agggacttga gtggatttga tggattttc ctggagaggg gagtaactgaa 180

tacaatgaga agttaaaggc cagggccaca ctgagtgtag acaagtccctc cagcacagcc 240

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cagtctccat cttcttggc tggatctcta gggcagaggg ccacccatcc ctgcagagcc 480

agtggaaatgt ttgataactta tggcgataact tttatgtact ggtaccagca gaaaccagga 540

cagccaccca aactcctcat ctatcttgc accaacctag gatctgggtt ccctgccagg 600

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gatgctgcaa cctattactg tcagcaaat aatgaggatc cgtacacgtt cggagggggc 720

accaagctgg aaatcaaactg tggtgaggc ggttcacccat gggcgccgc cttctccgtc 780

agtcccgct cgggtctgag tgacggacag agcgtgtcgg tgcgtcgat cgggtgccgc 840

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<211> 372
<212> PRT
<213> Chemically synthesized

<400> 2

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35 40 45

Ile Gly Trp Ile Phe Pro Gly Glu Gly Ser Thr Glu Tyr Asn Glu Lys
50 55 60

Phe Lys Gly Arg Ala Thr Leu Ser Val Asp Lys Ser Ser Ser Thr Ala
65 70 75 80

Tyr Met Glu Leu Thr Arg Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe
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Cys Ala Arg Gly Asp Tyr Tyr Arg Arg Tyr Phe Asp Leu Trp Gly Gln
100 105 110

Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Ser Gly Gly Gly
115 120 125

Gly Ser Asp Ile Glu Leu Ser Gly Gly Gly Thr Gln Ser Pro Ala
130 135 140

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145 150 155 160

Ser Glu Ser Val Asp Thr Tyr Gly Asp Thr Phe Met Tyr Trp Tyr Gln

165

170

175

Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Leu Ala Thr Asn
180 185 190

Leu Gly Ser Gly Val Pro Ala Gly Phe Ser Gly Ser Gly Ser Arg Thr
195 200 205

Asn Phe Thr Leu Thr Ile Asp Pro Val Glu Ala Asp Asp Ala Ala Thr
210 215 220

Tyr Tyr Cys Gln Gln Asn Asn Glu Asp Pro Tyr Thr Phe Gly Gly Gly
225 230 235 240

Thr Lys Leu Glu Ile Lys Arg Gly Gly Gly Ser Glu Phe Ala Pro
245 250 255

Ala Phe Ser Val Ser Pro Ala Ser Gly Leu Ser Asp Gly Gln Ser Val
260 265 270

Ser Val Ser Val Ser Gly Ala Ala Ala Gly Glu Thr Tyr Tyr Ile Ala
275 280 285

Gln Cys Ala Pro Val Gly Gly Gln Asp Ala Cys Asn Pro Ala Thr Ala
290 295 300

Thr Ser Phe Thr Thr Asp Ala Ser Gly Ala Ala Ser Phe Ser Phe Val
305 310 315 320

Val Arg Lys Ser Tyr Thr Gly Ser Thr Pro Glu Gly Thr Pro Val Gly
325 330 335

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